



contig name: [REDACTED]
contig assembler: jojeman
components:

[REDACTED]
[REDACTED]
[REDACTED] (99.3%)

assembly done: [REDACTED]

VECTOR_MATCH: NONE
VECTOR_MATCH: NONE
OLIGO_MATCH: NONE SEQ [REDACTED]

Length: 829

4:03 PM Check:

5459

POLYA_MATCH: MADE STRAND f

LOC e BEG 802 END 819 [REDACTED]

REMOVED f BEG 807 END 829

"AAAAAAAAAAAAACCGCGCCGC"

Length: 806

Check: 4601

REPMATCH: status:USEFUL 673 to 806 was Lib:B2 strand f

score 101 basecutoff 100 truecutoff 200 - NOT REPLACED with Ns

Length: 806

Check: 4601 ..

```
1  GGAACGAGGG AAAATCTGCC TTCTCACCAT GAGGCTTCTA GTCCTTTCCA
51 GCCTGCTCTG TATCCTGCTT CTCTGCTTCT CCATCTTCTC CACAGAAGGG
101 AAGAGGCGTC CTGCCAAGGC CTGGTCAGGC AGGAGAACCA GGCTCTGCTG
151 CCACCGAGTC CCTAGCCCCA ACTCAACAAA CCTGAAAGGA CATCATGTGA
201 GGCTCTGTAA ACCATGCAAG CTTGAGCCAG AGCCCCGCCT TTGGGTGGTG
251 CCTGGGGCAC TCCACAGGT GTAGCACTCC CAAAGCAAGA CTCCAGACAG
301 CGGAGAACCT CATGCCTGGC ACCTGAGGTA CCCAGCAGCC TCCTGTCTCC
351 CCTTTCAGCC TTCACAGCAG TGAGCTGCAA TGTTGGAGGG CTTTCATCTCG
401 GGCTGCAAGG ACCCTGGGAA AGTTCCAGAA CTCCACGTCC TTGTCTCAAT
451 TGTGCCATCA ACTTTCAGAG CTATCATGAG CCAACCTCAC CCCACAGGGC
501 CTCAGTCGCC ACCATGTGGG CCTCTCCAGT GCAAACCACC GAGCATTCCA
551 CCATGACCGG TCACAGCTAC AAATCCAGAG ACCATCAATC CTGCTAGAGT
601 GCAGGGTGGC AAGCACCCAA GGGTGGCTGA CCAAGACTGC AGAGTCTCCT
651 CCATCTTCAG GTCCATTGAG CCTCCTGGCA TTAACTACC AGCATCCAGT
701 GGTCCCCAAG GAATCCCTTC CTAGCCTCCT GACATGAGTC TGCTGGAAAG
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751 AGCATCCAAA CAAACAAGTA ATAAATAAAT AAATAAACTC AATGCAGACA

801 CAAAAA